SCORE Search Results Details for Application 09961086 and Search Result 20090302_142103_us-09-961-086a-1.rai.

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This page gives you Search Results detail for the Application 09961086 and Search Result 20090302_142103_us-09-961-086a-1.rai.

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OM protein - protein search, using sw model

Run on: March 3, 2009, 03:46:14; Search time 207 Seconds

(without alignments)

681.331 Million cell updates/sec

Title: US-09-961-086A-1

Perfect score: 3352

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Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 1316349 segs, 215321474 residues

Total number of hits satisfying chosen parameters: 1316349

Minimum DB seg length: 0

Maximum DB seg length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued_Patents_AA:*

1: /ABSS/Data/CRF/ptodata/1/iaa/5 COMB.pep:*

2: /ABSS/Data/CRF/ptodata/1/iaa/6_COMB.pep:*

3: /ABSS/Data/CRF/ptodata/1/iaa/7 COMB.pep:*

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7. /mbb/baca/ora/pedaaca/1/14a/na_oomb.pep.

7: /ABSS/Data/CRF/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed,

and is derived by analysis of the total score distribution.

SUMMARIES

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1	3352	100.0	655	2	US-09-245-808-1	Sequence 1, Appli
2	3346	99.8	655	3	US-11-333-542-6	Sequence 6, Appli
3	3342	99.7	655	3	US-11-333-542-8	Sequence 8, Appli
4	3331	99.4	655	2	US-09-767-594-1	Sequence 1, Appli
5	3331	99.4	655	2	US-09-584-586-10	Sequence 10, Appl
6 7	3331	99.4	655	3	US-09-856-927-2	Sequence 2, Appli
	3331	99.4	655	3	US-11-333-542-7	Sequence 7, Appli
8	3225	96.2	655	3	US-11-333-542-2	Sequence 2, Appli
9	3223.5	96.2	654	3	US-11-333-542-5	Sequence 5, Appli
10	2757	82.2	657	2	US-09-584-586-14	Sequence 14, Appl
11	835.5	24.9	1049	2	US-09-538-092-72	Sequence 72, Appl
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13	812	24.2	687	3	US-09-619-049-264	Sequence 264, App
14	795.5	23.7	676	3	US-10-369-493-3799	Sequence 3799, Ap
15	706.5	21.1	674	2	US-09-538-092-1125	Sequence 1125, Ap
16	702.5	21.0	663	3	US-10-473-696-6	Sequence 6, Appli
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33	623	18.6	147	2	US-09-584-586-12	Sequence 12, Appl
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35	612.5	18.3	1501	2	US-09-487-558B-346	Sequence 346, App
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39	594	17.7	1564	2	US-09-487-558B-244	Sequence 244, App
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41	589	17.6	1549	3	US-10-369-493-3919	Sequence 3919, Ap
42	580.5	17.3	1529	3	US-10-369-493-1692	Sequence 1692, Ap
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44	561.5	16.8	1395	3	US-10-369-493-4054	Sequence 4054, Ap
45	552.5	16.5	611	3	US-10-369-493-12397	Sequence 12397, A

ALIGNMENTS

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RESULT 1
US-09-245-808-1
; Sequence 1, Application US/09245808
; Patent No. 6313277
; GENERAL INFORMATION:
; APPLICANT: Doyle, L. Austin
  APPLICANT: Abruzzo, Lynne V.
  APPLICANT: Ross, Douglas D.
  TITLE OF INVENTION: Breast Cancer Resistance Protein (BCRP) and DNA which
  TITLE OF INVENTION: encodes it
  FILE REFERENCE: Ross UMb conversion
  CURRENT APPLICATION NUMBER: US/09/245,808
  CURRENT FILING DATE: 1999-02-05
  EARLIER APPLICATION NUMBER: 60/073763
  EARLIER FILING DATE: 1998-02-05
  NUMBER OF SEO ID NOS: 7
  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1
  LENGTH: 655
   TYPE: PRT
   ORGANISM: Human MCF-7/AdrVp cells
US-09-245-808-1
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  Best Local Similarity 100.0%; Pred. No. 0;
 Matches 655; Conservative 0; Mismatches 0; Indels 0; Gaps
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SCORE Search Results Details for Application 09961086 and Search Result 20090302_142103_us-09-961-086a-1.rai.
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US-11-333-542-6
; Sequence 6, Application US/11333542
; Patent No. 7465788
; GENERAL INFORMATION:
 APPLICANT: TAKEBE, NAOKO
  TITLE OF INVENTION: RHESUS BCRP AND ANTIBODIES THERETO
; FILE REFERENCE: UNIMD-0016
  CURRENT APPLICATION NUMBER: US/11/333,542
; CURRENT FILING DATE: 2006-01-18
  PRIOR APPLICATION NUMBER: 60/644,706
  PRIOR FILING DATE: 2005-01-18
 NUMBER OF SEO ID NOS: 13
  SOFTWARE: PatentIn Ver. 3.3
; SEO ID NO 6
  LENGTH: 655
   TYPE: PRT
  ORGANISM: Homo sapiens
US-11-333-542-6
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                          99.8%; Score 3346; DB 3; Length 655;
 Best Local Similarity 99.8%; Pred. No. 0;
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RESULT 3

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US-11-333-542-8

[;] Sequence 8, Application US/11333542

[;] Patent No. 7465788

[;] GENERAL INFORMATION:

[;] APPLICANT: TAKEBE, NAOKO

[;] TITLE OF INVENTION: RHESUS BCRP AND ANTIBODIES THERETO

[;] FILE REFERENCE: UNIMD-0016

[;] CURRENT APPLICATION NUMBER: US/11/333,542

[;] CURRENT FILING DATE: 2006-01-18

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PRIOR APPLICATION NUMBER: 60/644,706
  PRIOR FILING DATE: 2005-01-18
  NUMBER OF SEQ ID NOS: 13
  SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 8
  LENGTH: 655
   TYPE: PRT
   ORGANISM: Homo sapiens
US-11-333-542-8
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; Sequence 1, Application US/09767594
; Patent No. 6521635
; GENERAL INFORMATION:
 APPLICANT: Bates, Susan
; APPLICANT: Robey, Robert
  APPLICANT: The Government of the United States of America
  APPLICANT: as represented by the Secretary of the
  APPLICANT: Department of Health and Human Services
  TITLE OF INVENTION: Inhibition of MXR Transport by Acridine Derivatives
  FILE REFERENCE: 015280-402100US
  CURRENT APPLICATION NUMBER: US/09/767,594
  CURRENT FILING DATE: 2001-01-22
  PRIOR APPLICATION NUMBER: US 60/177,410
; PRIOR FILING DATE: 2000-01-20
  NUMBER OF SEO ID NOS: 2
  SOFTWARE: PatentIn Ver. 2.1
; SEO ID NO 1
  LENGTH: 655
   TYPE: PRT
  ORGANISM: Homo sapiens
  FEATURE:
  OTHER INFORMATION: human mitoxanthrone resistance (MXR)/BRCP/ABCP
   OTHER INFORMATION: protein
US-09-767-594-1
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SCORE Search Results Details for Application 09961086 and Search Result 20090302_142103_us-09-961-086a-1.rai.
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RESULT 5

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US-09-584-586-10
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; Sequence 10, Application US/09584586

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: Patent No. 6933150
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[;] GENERAL INFORMATION:

APPLICANT: Sorrentino, Brian

APPLICANT: Bunting, Kevin

TITLE OF INVENTION: EXPANSION OF HEMATOPOIETIC STEM CELLS TRANSDUCED WITH TITLE OF INVENTION: MDR-1 METHODS OF USE THEREOF

FILE REFERENCE: 1340-1-021CIP

CURRENT APPLICATION NUMBER: US/09/584,586

CURRENT FILING DATE: 2000-05-31

EARLIER APPLICATION NUMBER: US 60/086,988

EARLIER FILING DATE: 1998-05-28

EARLIER APPLICATION NUMBER: PCT/US99/11825

EARLIER FILING DATE: 1999-05-27

NUMBER OF SEO ID NOS: 16

SOFTWARE: PatentIn Ver. 2.0

[;] SEQ ID NO 10

LENGTH: 655

TYPE: PRT

; ORGANISM: Homo sapiens US-09-584-586-10

Ouerv Match 99.4%; Score 3331; DB 2; Length 655; Best Local Similarity 99.4%; Pred. No. 0; 0; Gaps Matches 651; Conservative 1; Mismatches 3: Indels 0; Qy 1 MSSSNVEVFIPVSOGNTNGFPATASNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE 60 Db 1 MSSSNVEVFIPVSOGNTNGFPATVSNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE 60 61 KEILSNINGIMKPGLNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGAPRPANFKCN 120 61 KEILSNINGIMKPGLNAILGPIGGGKSSLLDVLAARKDPSGLSGDVLINGAPRPANFKCN 120 Db Qy 121 SGYVVODDVVMGTLTVRENLOFSAALRLATTMTNHEKNERINRVIOELGLDKVADSKVGT 180 121 SGYVVODDVVMGTLTVRENLOFSAALRLATTMTNHEKNERINRVIEELGLDKVADSKVGT 180 181 OFIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMSKOGRTIIF 240 181 OFTRGVSGGERKRTSIGMELITDPSILSLDEPTTGLDSSTANAVLLLLKRMSKOGRTIIF 240 Db Qy 241 SIHOPRYSIFKLFDSLTLLASGRLMFHGPAOEALGYFESAGYHCEAYNNPADFFLDIING 300 241 SIHOPRYSIFKLFDSLTLLASGRLMFHGPAOEALGYFESAGYHCEAYNNPADFFLDIING 300 301 DSTAVALNREEDFKATEIIEPSKODKPLIEKLAEIYVNSSFYKETKAELHOLSGGEKKKK 360 301 DSTAVALNREEDFKATEIIEPSKQDKPLIEKLAEIYVNSSFYKETKAELHQLSGGEKKKK 360 Db Qу 361 ITVFKEISYTTSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVVLGLVIGAIYFGLKNDS 420 361 ITVFKEISYTTSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVVLGLVIGAIYFGLKNDS 420 Db 421 TGIQNRAGVLFFLTTNQCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDLLP 480 421 TGIONRAGVLFFLTTNOCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDLLP 480 Db Qy 481 MTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGOSVVSVATLL 540 481 MRMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGOSVVSVATLL 540 Qv 541 MTICEVEMMIESGLLVNLTTIASWLSWLOYFSIPRYGETALOHNEELGONECPGLNATGN 600 541 MTICFVFMMIFSGLLVNLTTIASWLSWLOYFSIPRYGFTALOHNEFLGONFCPGLNATGN 600 Db Qу 601 NPCNYATCTGEEYLVKOGIDLSPWGLWKNHVALACMIVIFLTIAYLKLLFLKKYS 655 601 NPCNYATCTGEEYLVKQGIDLSPWGLWKNHVALACMIVIFLTIAYLKLLFLKKYS 655 Db

RESHLT 6

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US-09-856-927-2
; Sequence 2, Application US/09856927
; Patent No. 7138493
: GENERAL INFORMATION:
; APPLICANT: Dean, Michael
; APPLICANT: Allikmets, Rando
  APPLICANT: Bates, Susan E.
  APPLICANT: Fojo, Antonio T.
  APPLICANT: The Government of the United States of America
  APPLICANT: as represented by the Secretary of the
  APPLICANT: Department of Health and Human Services
  TITLE OF INVENTION: A No. 7138493el ATP-Binding Cassette Protein Responsible for
  TITLE OF INVENTION: Cytotoxin Resistance
  FILE REFERENCE: 015280-382100US
  CURRENT APPLICATION NUMBER: US/09/856,927
  CURRENT FILING DATE: 2001-05-29
  PRIOR APPLICATION NUMBER: US 60/110,473
  PRIOR FILING DATE: 1998-11-30
  PRIOR APPLICATION NUMBER: WO PCT/US99/28107
  PRIOR FILING DATE: 1999-11-24
 NUMBER OF SEC ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
: SEO ID NO 2
  LENGTH: 655
   TYPE: PRT
  ORGANISM: Homo sapiens
US-09-856-927-2
 Query Match
                         99.4%; Score 3331; DB 3; Length 655;
 Best Local Similarity 99.4%; Pred. No. 0;
 Matches 651; Conservative 1; Mismatches 3; Indels 0; Gaps
                                                                          0;
           1 MSSSNVEVFIPVSOGNTNGFPATASNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE 60
0.v
Db
           1 MSSSNVEVFIPVSQGNTNGFPATVSNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE 60
Qv
          61 KEILSNINGIMKPGLNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGAPRPANFKCN 120
Db
          61 KEILSNINGIMKPGLNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGAPRPANFKCN 120
         121 SGYVVODDVVMGTLTVRENLOFSAALRLATTMTNHEKNERINRVIOELGLDKVADSKVGT 180
0v
Db
         121 SGYVVQDDVVMGTLTVRENLQFSAALRLATTMTNHEKNERINRVIEELGLDKVADSKVGT 180
         181 OFIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMSKOGRTIIF 240
QУ
Dh
         181 OFIRGVSGGERKRTSIGMELITDPSILSLDEPTTGLDSSTANAVLLLLKRMSKOGRTIIF 240
         241 SIHOPRYSIFKLFDSLTLLASGRLMFHGPAQEALGYFESAGYHCEAYNNPADFFLDIING 300
Q.V
Db
         241 SIHQPRYSIFKLFDSLTLLASGRLMFHGPAQEALGYFESAGYHCEAYNNPADFFLDIING 300
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301 DSTAVALNREEDFKATEIIEPSKODKPLIEKLAEIYVNSSFYKETKAELHOLSGGEKKKK 360
Qy
Dh
         301 DSTAVALNREEDFKATEIIEPSKODKPLIEKLAEIYVNSSFYKETKAELHOLSGGEKKKK 360
         361 ITVFKEISYTTSFCHQLRWVSKRSFKNLLGNPQASIAQIIVTVVLGLVIGAIYFGLKNDS 420
Db
         361 ITVFKEISYTTSFCHQLRWVSKRSFKNLLGNPQASIAQIIVTVVLGLVIGAIYFGLKNDS 420
         421 TGIONRAGVLFFLTTNOCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDLLP 480
QУ
         421 TGIONRAGVLFFLTTNOCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDLLP 480
Db
Qv
         481 MTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGQSVVSVATLL 540
Db
         481 MRMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGOSVVSVATLL 540
         541 MTICFVFMMIFSGLLVNLTTIASWLSWLOYFSIPRYGFTALOHNEFLGONFCPGLNATGN 600
QУ
Db
         541 MTICFVFMMIFSGLLVNLTTIASWLSWLOYFSIPRYGFTALOHNEFLGONFCPGLNATGN 600
         601 NPCNYATCTGEEYLVKOGIDLSPWGLWKNHVALACMIVIFLTIAYLKLLFLKKYS 655
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         601 NPCNYATCTGEEYLVKOGIDLSPWGLWKNHVALACMIVIFLTIAYLKLLFLKKYS 655
RESULT 7
US-11-333-542-7
; Sequence 7, Application US/11333542
; Patent No. 7465788
; GENERAL INFORMATION:
 APPLICANT: TAKEBE, NAOKO
  TITLE OF INVENTION: RHESUS BCRP AND ANTIBODIES THERETO
  FILE REFERENCE: UNIMD-0016
  CURRENT APPLICATION NUMBER: US/11/333,542
  CURRENT FILING DATE: 2006-01-18
  PRIOR APPLICATION NUMBER: 60/644,706
  PRIOR FILING DATE: 2005-01-18
  NUMBER OF SEQ ID NOS: 13
  SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 7
  LENGTH: 655
   TYPE: PRT
  ORGANISM: Homo sapiens
US-11-333-542-7
 Query Match
                         99.4%; Score 3331; DB 3; Length 655;
  Best Local Similarity 99.4%; Pred. No. 0;
 Matches 651; Conservative 1; Mismatches 3; Indels
                                                               0: Gaps
                                                                            0:
            1 MSSSNVEVFIPVSOGNTNGFPATASNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE 60
QУ
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US-11-333-542-2 ; Sequence 2, Application US/11333542 : Patent No. 7465788

; GENERAL INFORMATION: ; APPLICANT: TAKEBE, NAOKO

RESULT 8

; TITLE OF INVENTION: RHESUS BCRP AND ANTIBODIES THERETO

, TIBE OF INVENTION. MILEGO DOM THE INTEREST THE

; FILE REFERENCE: UNIMD-0016

CURRENT APPLICATION NUMBER: US/11/333,542

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CURRENT FILING DATE: 2006-01-18
; PRIOR APPLICATION NUMBER: 60/644,706
; PRIOR FILING DATE: 2005-01-18
 NUMBER OF SEO ID NOS: 13
  SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 2
  LENGTH: 655
   TYPE: PRT
   ORGANISM: Macaca mulatta
US-11-333-542-2
 Ouerv Match
                        96.2%; Score 3225; DB 3; Length 655;
 Best Local Similarity 96.2%; Pred. No. 0;
 Matches 630; Conservative 7; Mismatches 18; Indels
                                                           0; Gaps
                                                                       0;
Qy
           1 MSSSNVEVFIPVSOGNTNGFPATASNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE 60
             Db
           1 MSSSNVEVFIPMSOENTNGFPTTTSNDRKAFTEGAVLSFHNICYRVKVKSGFLPGRKPVE 60
          61 KEILSNINGIMKPGLNAILGPIGGKSSLLDVLAARKDPSGLSGDVLINGAPRPANEKCN 120
Db
          61 KEILSNINGIMKPGLNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGALRPTNFKCN 120
QУ
         121 SGYVVODDVVMGTLTVRENLOFSAALRLATTMTNHEKNERINRVIOELGLDKVADSKVGT 180
         121 SGYVVODDVVMGTLTVRENLOFSAALRLPTTMTNHEKNERINRVIOELGLDKVADSKVGT 180
Dh
         181 OFIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMSKOGRTIIF 240
         181 QFIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMSKQGRTIIF 240
Db
QУ
         241 SIHOPRYSIFKLFDSLTLLASGRLMFHGPAOEALGYFESAGYHCEAYNNPADFFLDIING 300
Db
         241 STHOPRYSIFKLFDSLTLLASGRLMFHGPAOEALGYFESAGYHCEAYNNPADFFLDIING 300
         301 DSTAVALNREEDFKATEIIEPSKQDKPLIEKLAEIYVNSSFYKETKAELHQLSGGEKKKK 360
         301 DSTAVALNREEDFKATEIIEPSKRDKPLVEKLAEIYVDSPFYKETKAELHOLSGGEKKKK 360
Db
QУ
         361 ITVFKEISYTTSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVVLGLVIGAIYFGLKNDS 420
         361 ITVFKEISYTTSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVILGLVIGGIYFGLNNDS 420
Dh
Qv
         421 TGIQNRAGVLFFLTTNQCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDLLP 480
         421 TGIONRAGVLFFLTTNOCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFFGKLLSDLLP 480
Db
QУ
         481 MTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGOSVVSVATLL 540
             Db
         481 MRMLPSIIFTCIVYFMLGLKPTADAFFIMMFTLMMVAYSASSMALAIAAGOSVVSVATLL 540
         541 MTICFVFMMIFSGLLVNLTTIASWLSWLQYFSIPRYGFTALQHNEFLGQNFCPGLNATGN 600
Qy
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241 SIHOPRYSIFKLFDSLTLLASGRLMFHGPAOEALGYFESAGYHCEAYNNPADFFLDIING 300

Dh

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Qу
         301 DSTAVALNREEDFKATEIIEPSKODKPLIEKLAEIYVNSSFYKETKAELHOLSGGEKKKK 360
Db
         301 DSTAVALNREEDFKATEIIEPSKRDKPLVEKLAEIYVDSSFYKETKAELHQLSGGE-KKK 359
         361 ITVFKEISYTTSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVVLGLVIGAIYFGLKNDS 420
QУ
Db
         360 ITVFKEISYTTSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVILGLVIGAIYFGLNNDS 419
         421 TGIONRAGVLFFLTTNOCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDLLP 480
Db
         420 TGIQNRAGVLFFLTTNQCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFFGKLLSDLLP 479
         481 MTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGOSVVSVATLL 540
QУ
         480 MRMLPSIIFTCIVYFMLGLKPTADAFFIMMFTLMMVAYSASSMALAIAAGOSVVSVATLL 539
Db
Qv.
         541 MTICFVFMMIFSGLLVNLTTIASWLSWLQYFSIPRYGFTALQHNEFLGQNFCPGLNATGN 600
         540 MTICFVFMMIFSGLLVNLTTIASWLSWLOYFSIPRYGFTALOHNEFLGONFCPGLNATVN 599
Db
         601 NPCNYATCTGEEYLVKOGIDLSPWGLWKNHVALACMIVIFLTIAYLKLLFLKKYS 655
QУ
             Db
         600 NTCNYATCTGEEYLAKOGIDLSPWGLWKNHVALACMIVIFLTIAYLKLLFLKKYS 654
RESHLT 10
US-09-584-586-14
; Sequence 14, Application US/09584586
; Patent No. 6933150
; GENERAL INFORMATION:
; APPLICANT: Sorrentino, Brian
; APPLICANT: Bunting, Kevin
  TITLE OF INVENTION: EXPANSION OF HEMATOPOIETIC STEM CELLS TRANSDUCED WITH
  TITLE OF INVENTION: MDR-1 METHODS OF USE THEREOF
  FILE REFERENCE: 1340-1-021CIP
  CURRENT APPLICATION NUMBER: US/09/584,586
  CURRENT FILING DATE: 2000-05-31
  EARLIER APPLICATION NUMBER: US 60/086,988
  EARLIER FILING DATE: 1998-05-28
  EARLIER APPLICATION NUMBER: PCT/US99/11825
  EARLIER FILING DATE: 1999-05-27
  NUMBER OF SEO ID NOS: 16
  SOFTWARE: PatentIn Ver. 2.0
: SEO ID NO 14
  LENGTH: 657
   TYPE: PRT
; ORGANISM: Mus musculus
US-09-584-586-14
                        82.2%; Score 2757; DB 2; Length 657;
 Query Match
  Best Local Similarity 81.5%; Pred. No. 3.4e-278;
 Matches 536; Conservative 51; Mismatches 67; Indels
                                                             4; Gaps
                                                                          3;
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Qу	1	MSSSNVEVFIPVSQGNTNGFPATASNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE	60
Db	1	MSSSNDHVLVPMSQRNNNGLPRMNSRAVRTLAEGDVLSFHHITYRVKVKSGFL-VRKTVE	59
Qу	61	KEILSNINGIMKPGLNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGAPRPANFKCN	120
Db	60	KEILSDINGIMKPGLNAILGPTGGGKSSLLDVLAARKDPKGLSGDVLINGAPQPAHFKCC	119
Qу	121	SGYVVQDDVVMGTLIVRENLQFSAALRLATIMINHEKNERINRVIQELGLDKVADSKVGT	180
Db	120	${\tt SGYVVQDDVVMGTLTVRENLQFSAALRLPTTMKNHEKNERINTIIKELGLEKVADSKVGT}$	179
QУ	181	QFIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMSKQGRTIIF	240
Db	180	${\tt QFIRGISGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMSKQGRTIIF}$	239
QУ	241	SIHQPRYSIFKLFDSLTLLASGRLMFHGPAQEALGYFESAGYHCEAYNNPADFFLDIING	300
Db	240	$\verb SIHQPRYSIFKLFDSLTLLASGKLVFHGPAQKALEYFASAGYHCEPYNNPADFFLDVING \\$	299
QУ	301	DSTAVALNREE-DFKATEIIEPSKQDKPLIEKLAEIYVNSSFYKETKAELHQLSGGEKKK	359
Db	300	${\tt DSSAVMLNREEQDNEANKTEEPSKGEKPVIENLSEFYINSAIYGETKAELDQLPGAQEKK}$	359
Qу	360	KITVFKEISYTTSFCHQLRWVSKRSFKNLLGNPQASIAQIIVTVVLGLVIGAIYFGLKND :	419
Db	360	GTSAFKEPVYVTSFCHQLRWIARRSFKNLLGNPQASVAQLIVTVILGLIIGAIYFDLKYD	419
Qу	420	STGIQNRAGVLFFLTTNQCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDLL: :	479
Db	420	${\tt AAGMQNRAGVLFFLTINQCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFFGKVMSDLL}$	479
Qу		PMTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGQSVVSVATL	
Db	480	${\tt PMRFLPSVIFTCILYFMLGLKKTVDAFFIMMFTLIMVAYTASSMALAIATGQSVVSVATL}$	539
Qу		LMTICFVFMMIFSGLLVNLTTIASWLSWLQYFSIPRYGFTALQHNEFLGQNFCPGLNATG	
Db		$\verb LMTIAFVFMMLFSGLLVNLRTIGPWLSWLQYFSIPRYGFTALQYNEFLGQEFCPGFNVTD \\$	
Qy		NNPCNYATCTGEEYLVKQGIDLSPWGLWKNHVALACMIVIFLTIAYLKLLFLKKYS 6	
Db	600	NSTCVNSYAICTGNEYLINQGIELSPWGLWKNHVALACMIIIFLTIAYLKLLFLKKYS 6	57

RESULT 11 US-09-538-092-72

; Sequence 72, Application US/09538092

; Patent No. 6753314 ; GENERAL INFORMATION: ; APPLICANT: Giot, Loic

```
; APPLICANT: Mansfield, Traci A.
: TITLE OF INVENTION: Protein-Protein Complexes and Method of Using Same
; FILE REFERENCE: 15966-542
; CURRENT APPLICATION NUMBER: US/09/538,092
; CURRENT FILING DATE: 2000-03-29
; PRIOR APPLICATION NUMBER: 60/127,352
 PRIOR FILING DATE: 1999-04-01
; PRIOR APPLICATION NUMBER: 60/178,965
; PRIOR FILING DATE: 2000-02-01
; NUMBER OF SEO ID NOS: 1387
 SOFTWARE: CuraPatSegFormatter Version 0.9
; SEQ ID NO 72
  LENGTH: 1049
  TYPE: PRT
 ORGANISM: Saccharomyces cerevisiae
 FEATURE:
; NAME/KEY: misc feature
; LOCATION: (0)...(0)
; OTHER INFORMATION: Polypeptide Accession Number YCR011C
US-09-538-092-72
                   24.9%; Score 835.5; DB 2; Length 1049;
 Query Match
 Best Local Similarity 30.5%; Pred. No. 4.7e-77;
 Matches 222; Conservative 134; Mismatches 257; Indels 115; Gaps 18;
         1 MSSSNVEVFIPVSOGNTNGFPATASNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE 60
QУ
           Db
       355 LGSSKSPIRLP-DEDAVNNFLQNEDDTL-----ATLSFENITYSVPSINS-----DGVE 402
       61 KEILSNINGIMKPG-LNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGAPRP-ANFK 118
Qv
           Db
       403 ETVLNEISGIVKPGOILAIMGGSGAGKTTLLDILAMKRKTGHVSGSIKVNGISMDRKSFS 462
       119 CNSGYVVODDVVMGTLTVRENLOFSAALRLATTMTNHEKNERINRVIOELGLDKVADSKV 178
0v
             Db
       463 KIIGFVDQDDFLLPTLTVFETVLNSALLRLPKALSFEAKKARVYKVLEELRIIDIKDRII 522
Qv
       179 GTOFIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMSKO-GRT 237
           Db
       523 GNEFDRGISGGEKRRVSIACELVTSPLVLFLDEPTSGLDASNANNVIECLVRLSSDYNRT 582
       238 IIFSIHOPRYSIFKLFDSLTLLASGRLMFHGPAOEALGYFESAGYHCEAYNNPADFFLDI 297
Qу
           Db
       583 LVLSTHOPRSNIFYLFDKLVLLSKGEMVYSGNAKKVSEFLRNEGYTCPDNYNTADYLIDI 642
QУ
       298 -----INGDSTAV 305
Dh
       643 TFEAGPOGKRRRIRNISDLEAGTDTNDIDNTIHOTTFTSSDGTTOREWAHLAAHRDEIRS 702
0.v
       306 ALNREEDFKATE----IIEPSKODKPLIEKLAEIYVNSSFYKETKAELHO-LSGGEKKKK 360
           Db
       703 LLRDEEDVEGTDGRRGATEIDLNTKLLHDK----YKDSVYYAELSQEIEEVLSEGDEESN 758
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361 IT--VFKEISYTTSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVVLGLVIGAIYFGLKN 418
Qy
                 : | || :: || || :: |
Dh
        759 VLNGDLPTGOOSAGFLOOLSILNSRSFKNMYRNPKLLLGNYLLTILLSLFLGTLYYNVSN 818
        419 DSTGIONRAGVLFFLTTNOCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDL 478
0.7
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        479 LPMTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGOSVVSVAT 538
QУ
            :|: ::| |: : ||| | | | | | | | :||| : |:: | | | | | | | :: :
Db
        879 VPLRVVPPILLSLIVYPMTGLNMKDNAFFKCIGILILFNLGISLEILTIGIIFEDLNNSI 938
        539 LLMTICFVFMMIFSGLLV---NLTTIASWLSWLQYFSIPRYGFTALQHNEF---- 586
Qv
            :| : : ::|||| : |:| :| :|: ||: | : :| ||
        939 ILSVLVLLGSLLFSGLFINTKNITNVA--FKYLKNFSVFYYAYESLLINEVKTLMLKERK 996
Db
       587 LGONF-CPGLNATGNNPCNYATCTGEEYLVKOGI--DLSPWGLWKNHVALACMIVIFLTI 643
QУ
            997 YGLNIEVPG-----ATILSTFGFVVONLVFDIK-----ILALFNVVFLIM 1036
Db
       644 AYLKLLFL 651
Qv
            11 1 ::
Db 1037 GYLALKWI 1044
RESULT 12
US-10-369-493-1520
; Sequence 1520, Application US/10369493
: Patent No. 7314974
; GENERAL INFORMATION:
; APPLICANT: Cao, Yongwei
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; APPLICANT: Goldman, Barry S.
; APPLICANT: Chen, Xianfeng
 TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
; TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
; FILE REFERENCE: 38-10(52052)B
; CURRENT APPLICATION NUMBER: US/10/369,493
; CURRENT FILING DATE: 2003-02-28
; PRIOR APPLICATION NUMBER: US 60/360,039
; PRIOR FILING DATE: 2002-02-21
; NUMBER OF SEQ ID NOS: 47374
; SEO ID NO 1520
  LENGTH: 1049
; TYPE: PRT
; ORGANISM: Saccharomyces cerevisiae
US-10-369-493-1520
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http://es/ScoreAccessWeb/GetItem.action?AppId=099610...2 142103 us-09-961-086a-1.rai&ItemType=4&startByte=0 (18 of 24)3/11/2009 3:07:47 PM

Query Match 24.9%; Score 835.5; DB 3; Length 1049;

Best Local Similarity 30.5%; Pred. No. 4.7e-77;

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Matches 222; Conservative 134; Mismatches 257; Indels 115; Gaps 18;
         1 MSSSNVEVFIPVSQGNTNGFPATASNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVE 60
QУ
          Db
       355 LGSSKSPIRLP-DEDAVNNFLQNEDDTL-----ATLSFENITYSVPSINS-----DGVE 402
       61 KEILSNINGIMKPG-LNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGAPRP-ANFK 118
Qу
          : :|: |:||:|| : ||:| :| ||:||:|| :: ||:|| :: ||:||
Db
       403 ETVLNEISGIVKPGOILAIMGGSGAGKTTLLDILAMKRKTGHVSGSIKVNGISMDRKSFS 462
       119 CNSGYVVQDDVVMGTLTVRENLQFSAALRLATTMTNHEKNERINRVIQELGLDKVADSKV 178
Qy
             463 KIIGFVDODDFLLPTLTVFETVLNSALLRLPKALSFEAKKARVYKVLEELRIIDIKDRII 522
Db
Qу
       179 GTOFIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLKRMSKO-GRT 237
          Db
       523 GNEFDRGISGGEKRRVSIACELVTSPLVLFLDEPTSGLDASNANNVIECLVRLSSDYNRT 582
       238 IIFSIHOPRYSIFKLFDSLTLLASGRLMFHGPAOEALGYFESAGYHCEAYNNPADFFLDI 297
Qy
          583 LVLSIHOPRSNIFYLFDKLVLLSKGEMVYSGNAKKVSEFLRNEGYICPDNYNIADYLIDI 642
Db
       298 -----INGDSTAV 305
Qу
Db
       643 TFEAGPQGKRRRIRNISDLEAGTDTNDIDNTIHQTTFTSSDGTTQREWAHLAAHRDEIRS 702
       306 ALNREEDFKATE----IIEPSKODKPLIEKLAEIYVNSSFYKETKAELHO-LSGGEKKKK 360
Qy
           703 LLRDEEDVEGTDGRRGATEIDLNTKLLHDK----YKDSVYYAELSOEIEEVLSEGDEESN 758
Db
       361 IT--VFKEISYTTSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVVLGLVIGAIYFGLKN 418
                  : | || :: || ||: : :: |:: | : | : |
       759 VLNGDLPTGQQSAGFLQQLSILNSRSFKNMYRNPKLLLGNYLLTILLSLFLGTLYYNVSN 818
QУ
       419 DSTGIONRAGVLFFLTTNOCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDL 478
          819 DISGFONRMGLFFFILTYFGFVTFTGLSSFALERIIFIKERSNNYYSPLAYYISKIMSEV 878
Db
       479 LPMTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGQSVVSVAT 538
QУ
          879 VPLRVVPPILLSLIVYPMTGLNMKDNAFFKCIGILILFNLGISLEILTIGIIFEDLNNSI 938
Db
       539 LLMTICFVFMMIFSGLLV---NLTTIASWLSWLQYFSIPRYGFTALQHNEF----- 586
QУ
          :| : : ::|||| : |:||:| :|: ||: ||: ||
       939 ILSVLVLLGSLLFSGLFINTKNITNVA--FKYLKNFSVFYYAYESLLINEVKTLMLKERK 996
Db
       587 LGONF-CPGLNATGNNPCNYATCTGEEYLVKQGI--DLSPWGLWKNHVALACMIVIFLTI 643
          997 YGLNIEVPG-----ATILSTFGFVVQNLVFDIK-----ILALFNVVFLIM 1036
Db
       644 AYLKLLFL 651
QУ
           11 1 ::
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Db 1037 GYLALKWI 1044
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RESHLT 13
US-09-619-049-264
; Sequence 264, Application US/09619049
; Patent No. 7135558
: GENERAL INFORMATION:
; APPLICANT: YANDELL, MARK
 TITLE OF INVENTION: ISOLATED DROSOPHILA PROTEINS ESSENTIAL
; TITLE OF INVENTION: FOR SURVIVAL, NUCLEIC ACID MOLECULES ENCODING ESSENTIAL
; TITLE OF INVENTION: DROSOPHILA PROTEINS, AND USES THEREOF AS INSECTICIDAL
  TITLE OF INVENTION: TARGETS
; FILE REFERENCE: CL000735
; CURRENT APPLICATION NUMBER: US/09/619,049
 CURRENT FILING DATE: 2000-07-18
; PRIOR APPLICATION NUMBER: 60/171,590
; PRIOR FILING DATE: 1999-12-23
  PRIOR APPLICATION NUMBER: 60/171,627
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: 60/175,763
 PRIOR FILING DATE: 2000-01-12
; PRIOR APPLICATION NUMBER: 60/175,685
; PRIOR FILING DATE: 2000-01-12
; PRIOR APPLICATION NUMBER: 60/186,663
; PRIOR FILING DATE: 2000-03-03
: PRIOR APPLICATION NUMBER: 60/187,241
; PRIOR FILING DATE: 2000-03-03
; NUMBER OF SEO ID NOS: 1533
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEO ID NO 264
; LENGTH: 687
; TYPE: PRT
  ORGANISM: DROSOPHILA
US-09-619-049-264
 Ouerv Match
                      24.2%; Score 812; DB 3; Length 687;
 Best Local Similarity 32.1%; Pred. No. 6.6e-75;
 Matches 210; Conservative 134; Mismatches 251; Indels 60; Gaps 17;
         5 NVEVFIPVSOGNTNGFPATASNDLKAFTEGAVLSFHNICYRVKLKSGFLPCRKPVEKEIL 64
Qv
            Db
         74 NMDIFGAVNO-----PGSGWROLVNRTRGLFCNERHI-----PAPR---KHLL 113
        65 SNINGIMKPG-LNAILGPTGGGKSSLLDVLAARKDPSGL---SGDVLINGAPRPA-NFK 118
Ov
             |: |: || | |::| :| ||::||: || | | : || :|| | : ||
Db
     114 KNVCGVAYPGELLAVMGSSGAGKTTLLNALAFR-SPOGIOVSPSGMRLLNGOPVDAKEMO 172
       119 CNSGYVVQDDVVMGTLTVRENLQFSAALRLATTMTNHEKNERINRVIQELGLDKVADSKV 178
Qv
                Db 173 ARCAYVOODDLFIGSLTAREHLIFOAMVRMPRHLTYRORVARVDOVIOELSLSKCOHTII 232
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Qу
       179 GTO-FIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMSKOGRT 237
              Db
       233 GVPGRVKGLSGGERKRLAFASEALTDPPLLICDEPTSGLDSFTAHSVVQVLKKLSQKGKT 292
       238 IIFSIHOPRYSIFKLFDSLTLLASGRLMFHGPAOEALGYFESAGYHCEAYNNPADFFLDI 297
Qy
          Db
       293 VILTIHOPSSELFELFDKILLMAEGRVAFLGTPSEAVDFFSYVGAOCPTNYNPADFYVOV 352
0.v
       298 INGDSTAVALNREEDFKATEIIEPSKODKPLIEKLAEIYVNSSFYKETKAELHOLSGGEK 357
             353 L----AVVPGREIESR-----DRIAKICDNFAISKVAR-DMEQLLATKN 391
Db
       358 KKKITVFKEISYT--TSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVVLGLVIGAIYFG 415
Qy
           392 LEKPLEOPENGYTYKATWFMOFRAVLWRSWLSVLKEPLLVKVRLIOTTMVAILIGLIFLG 451
Db
0.v
       416 LKNDSTGIQNRAGVLFFLTTNQCFSSVSA-VELFVVEKKLFIHEYISGYYRVSSYFLGKL 474
           452 OOLTOVGVMNINGAIFLFLTNMTFONVFATINVFTSELPVFMREARSRLYRCDTYFLGKT 511
Db
       475 LSDLLPMTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGOSVV 534
QУ
          ::: ||: : ::|| | | |:||: || | |:|| ::| || ::|
Db
       512 IAE-LPLFLTVPLVFTAIAYPMIGLRAGVLHFFNCLALVTLVANVSTSFGYLISCASSST 570
Qv
       535 SVATLLMTICFVFMMIFSGLLVNLTTIASWLSWLQYFSIPRYGFTALQHNEFLGQNFCPG 594
          571 SMALSVGPPVIIPFLLFGGFFLNSGSVPVYLKWLSYLSWFRYANEGLLINOWADVE--PG 628
Db
      595 -LNATGNNPCNYATCTGEEYLVKOGIDLSPWGLWKNHVALACMIVIFLTIAYLKL 648
Qy
           Db
      629 EISCTSSN----TTCPSSGKVILETLNFSAADLPLDYVGLAILIVSFRVLAYLAL 679
RESULT 14
US-10-369-493-3799
; Sequence 3799, Application US/10369493
; Patent No. 7314974
: GENERAL INFORMATION:
```

```
; APPLICANT: Cao, Yongwei
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
 APPLICANT: Goldman, Barry S.
  APPLICANT: Chen, Xianfeng
  TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
  TITLE OF INVENTION: PLANTS WITH IMPROVED PROPERTIES
; FILE REFERENCE: 38-10(52052)B
; CURRENT APPLICATION NUMBER: US/10/369,493
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; CURRENT FILING DATE: 2003-02-28 : PRIOR APPLICATION NUMBER: US 60/360,039

; PRIOR FILING DATE: 2002-02-21

; NUMBER OF SEQ ID NOS: 47374 ; SEQ ID NO 3799

; LENGTH: 676 ; TYPE: PRT

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ORGANISM: Neurospora crassa
IIS-10-369-493-3799
                  23.7%; Score 795.5; DB 3; Length 676;
 Query Match
 Best Local Similarity 31.2%; Pred. No. 3.4e-73;
 Matches 199; Conservative 107; Mismatches 218; Indels 113; Gaps 11;
       61 KEILSNINGIMKPG-LNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGAP-RPANFK 118
QУ
          1 KEILSGIOGMAHPGEVTAIMGASGAGKTTFLDILARKNKRGOVSGDFYINGEKVSDPEYK 60
Db
Qv
    119 CNSGYVVQDDVVMGTLTVRENLQFSAALRLATTMTNHEKNERINRVIQELGLDKVADSKV 178
            Db
       61 NAVGFVDOEDTMLPTLTVHETILNSALLRLPKDMTRAAKEORVIEVEKOLGIYHIRDSLI 120
       179 GTO--FIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTA-NAVLLLLKRMSKOG 235
QУ
          Db
       121 GSEEGKGRGISGGEKRRYGIACELVISPSILFLDEPISGLDAYNAYNVVECLVILAKTYK 180
       236 RTIIFSIHQPRYSIFKLFDSLTLLASGRLMFHGPAQEALGYFESAGYHCEAYNNPADFFL 295
Qv
          Db
       181 RTVIFTIHQPRSNIVALFDRLILLAQGKTVYSGPLHQCQEYFDQIGYTCPPGFNIADYLV 240
       296 DI-----TEJIEPS----- 322
Qу
                     :: | :| : |:
Db
       241 DLTMHAGSTSSYDDGTLSVDGVSVGPSSTRAVKSIASVSGVSIGDDSLVESSSSRPRNKR 300
       323 -----KODKPL----- 328
Qv
              :1:: 1
Db
       301 RDSVRRROERELYTRRKOAVDTAASSDAGDEIGGYKLOKOPPVTPLRSTNDDLHDLPPLA 360
       329 ----IEKLAEIYVNSSFYKETKAELHOL-----SGGEKKKKITVFKEISYT----- 370
0v
              :: | | |::| | |:|| | | |:
Db
       361 ATGTDLDVLIESYIHSDIAASTHEEIHQAIAAAVNSNGQNSNGYVADGNI-YTGTMGKGY 419
Qv
       371 --TSFCHOLRWVSKRSFKNLLGNPOASIAOIIVTVVLGLVIGAIYFGLKNDSTGIONRAG 428
               420 ARVGLFRQFVILSQRTWKNLYRNPMLMLTHYAIAILLAVFAGYLFYGLTLDIAGFQNRLG 479
Db
       429 VLFFLTTNOCFSSVSAVELFVVEKKLFIHEYISGYYRVSSYFLGKLLSDLLPMTMLPSII 488
Qу
          Db
       480 LFFFVLALFGFSTLTSLGVFSQERLLFVRERANGYYSPITYFAAKVLFDIVPLRIIPPIL 539
       489 FTCIVYFMLGLKPKADAFFVMMFTLMMVAYSASSMALAIAAGOSVVSVATLLMTICFVFM 548
QУ
            Db
       540 LGAIIYPMTGLVADYORFFVFILVLVLFNLAAAAICLFIGILCKDGGVANLIGSLVMLFS 599
       549 MIFSGLLVNLTTIASWLSWLQYFSIPRYGFTALQHNE 585
0.v
          600 LLFAGLLLNHNAIPAAALWLQWLSIFHYGFEALIVNE 636
Db
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RESULT 15
US-09-538-092-1125
; Sequence 1125, Application US/09538092
; Patent No. 6753314
; GENERAL INFORMATION:
; APPLICANT: Giot, Loic
; APPLICANT: Mansfield, Traci A.
 TITLE OF INVENTION: Protein-Protein Complexes and Method of Using Same
; FILE REFERENCE: 15966-542
; CURRENT APPLICATION NUMBER: US/09/538,092
; CURRENT FILING DATE: 2000-03-29
; PRIOR APPLICATION NUMBER: 60/127,352
; PRIOR FILING DATE: 1999-04-01
  PRIOR APPLICATION NUMBER: 60/178,965
; PRIOR FILING DATE: 2000-02-01
; NUMBER OF SEO ID NOS: 1387
 SOFTWARE: CuraPatSegFormatter Version 0.9
; SEO ID NO 1125
; LENGTH: 674
  TYPE: PRT
 ORGANISM: Homo sapiens
: FEATURE:
 NAME/KEY: misc_feature
; LOCATION: (0) ... (0)
; OTHER INFORMATION: Polypeptide Accession Number P45844
US-09-538-092-1125
 Query Match
              21.1%; Score 706.5; DB 2; Length 674;
 Best Local Similarity 28.4%; Pred. No. 6.8e-64;
 Matches 194; Conservative 155; Mismatches 251; Indels 83; Gaps 23;
         3 SSNVEVFIPVSOGNTNGFPATASNDL---KAFT----EGAV-LSFHNICYRVKLKSGFLP 54
0.v
           34 SSNMEA---TETDLLNGHLKKVDNNLTEAQRFSSLPRRAAVNIEFRDLSYSVPEGPWW-- 88
Db
        55 CRKPVEKEILSNINGIMKPG-LNAILGPTGGGKSSLLDVLAARKDPSGLSGDVLINGAPR 113
Qv
            Db
        89 -RKKGYKTLLKGISGKFNSGELVAIMGPSGAGKSTLMNILAGYRE-TGMKGAVLINGLPR 146
       114 PAN-FKCNSGYVVODDVVMGTLTVRENLOFSAALRLATTMTNHEKNERINRVIOELGLDK 172
0v
               |: | |::|||::: |||:| : || |:| : : | : :: |||
Db
        147 DLRCFRKVSCYIMQDDMLLPHLTVQEAMMVSAHLKLQE--KDEGRREMVKEILTALGLLS 204
        173 VADSKVGTOFIRGVSGGERKRTSIGMELITDPSILFLDEPTTGLDSSTANAVLLLLKRMS 232
Qy
             Dh
        205 CANTRTGS----LSGGQRKRLAIALELVNNPPVMFFDEPTSGLDSASCFQVVSLMKGLA 259
0.v
        233 KOGRTIIFSIHOPRYSIFKLFDSLTLLASGRLMFHGPAQEALGYFESAGYHCEAYNNPAD 292
            Db
        260 QGGRSIICTIHQPSAKLFELFDQLYVLSQGQCVYRGKVCNLVPYLRDLGLNCPTYHNPAD 319
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Qу	293	FFLDIINGDSTAVALNREEDFKATEIIEPSKQDKPLIEKLAEIYVNSSFYKETKAEL-	349
Db	320	FVMEVASGEYGDQNSRLVRAVREGMCDSDHKRDLGGDAEVN	360
QУ	350	HQLSGGEKK-KKITVFKEISYTTSFCHQLRWVSKRSFKNLLGNPQASI	396
Db	361	${\tt PFLWHRPSEEVKQTKRLKGLRKDSSSMEGCHSFSASCLTQFCILFKRTFLSIMRDSVLTH}$	420
QУ	397	AQIIVTVVLGLVIGAIYFGLKNDSTGIQNRAGVLFFLTTNQCFSSVSAVEL-FVVEKKLF : :: : : : : : : : : :	455
Db	421	$\tt LRITSHIGIGLLIGLLYLGIGNEAKKVLSNSGFLFFSMLFLMFAALMPTVLTFPLEMGVF$	480
ДÄ	456	IHEYISGYYRVSSYFLGKLLSDLLPMTMLPSIIFTCIVYFMLGLKPKADAFFVMMFTL-M : ::::: :: :: : : : : : : : : : : :	514
Db	481	$\tt LREHLNYWYSLKAYYLAKTMAD-VPFQIMFPVAYCSIVYWMTS-QPSDAVRFVLFAALGT$	538
QУ	515	MVAYSASSMALAIAAGQSVVSVATLLMTICFVFMMIFSGLLVNLTTIASWLSWLQYFSIP : : : : : : : : : : : : : :	574
Db	539	$\verb MTSLVAQSLGLLIGAASTSLQVATFVGPVTAIPVLLFSGFFVSFDTIPTYLQWMSYISYV $	598
QУ	575	RYGFTALQHNEFLGQNFCPGLNATGNNPCNYATCTGEEYLVKQGIDLSPWGLW	627
Db	599	RYGFEGVILSIYGLDREDLHCDIDETCHFQKSEAILRE-LDVENAKLY	645
QУ	628	KNHVALACMIVIFLTIAYLKLLF 650 :: : :	
Db	646	LDFIVLGIFFISLRLIAYFVLRY 668	

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